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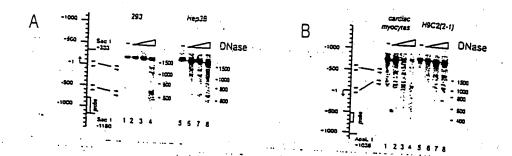
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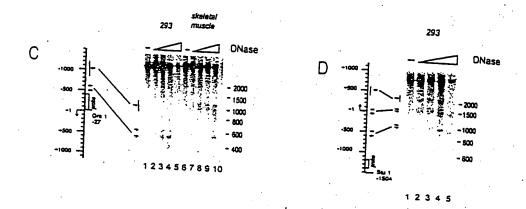
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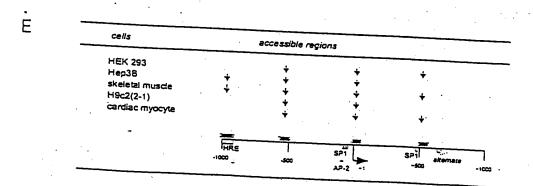
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Figure 1







Sequence conservation:

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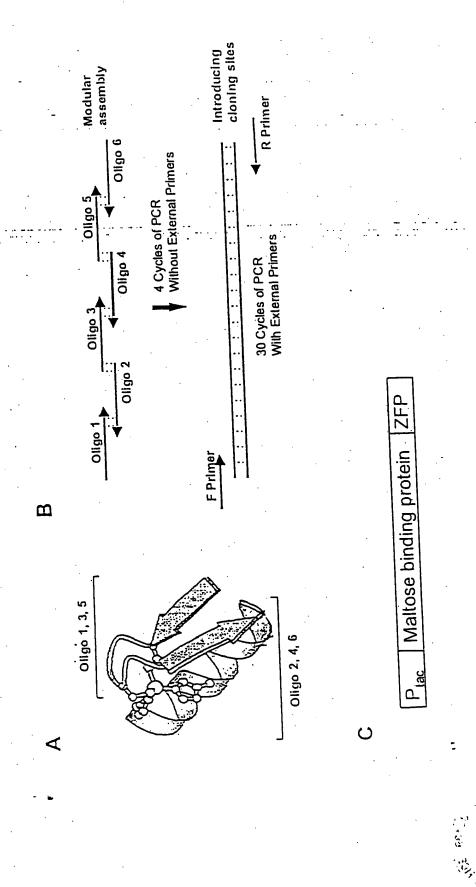


Figure 2

Figure 2

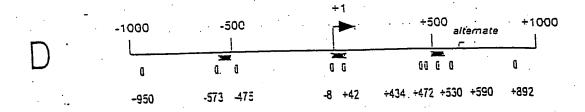
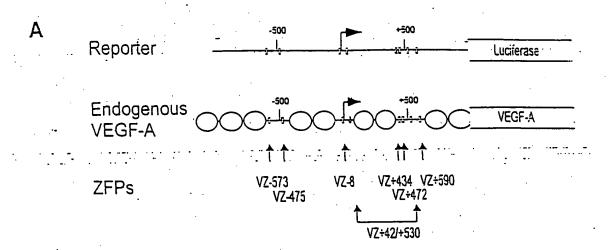


Figure 3



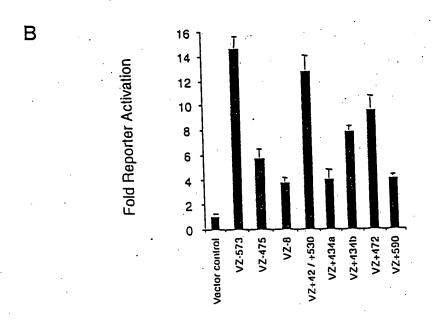
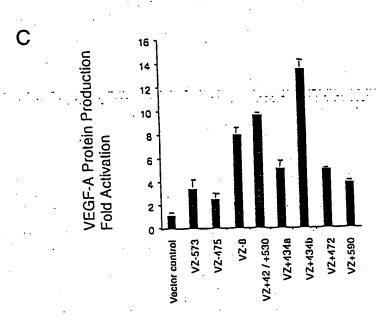


Figure 3



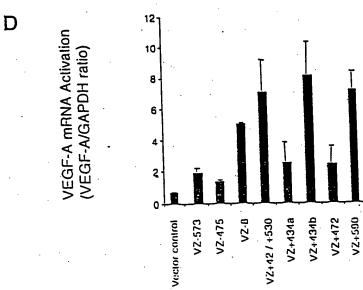


Figure 3

E

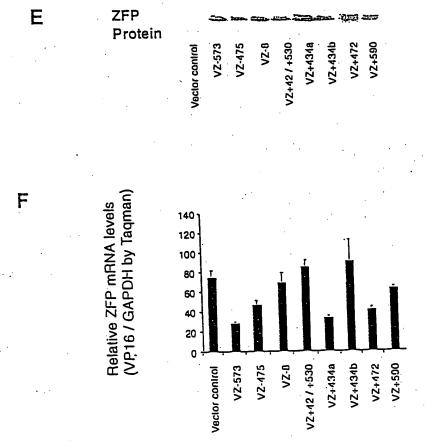
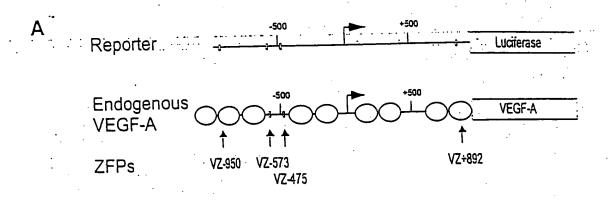
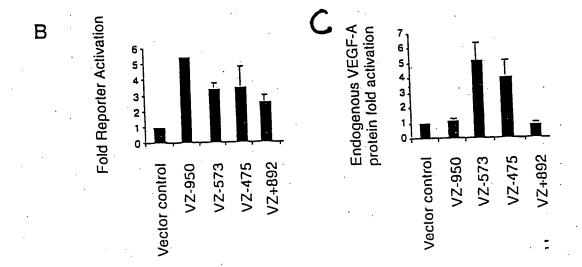


Figure 4





1

Figure 5

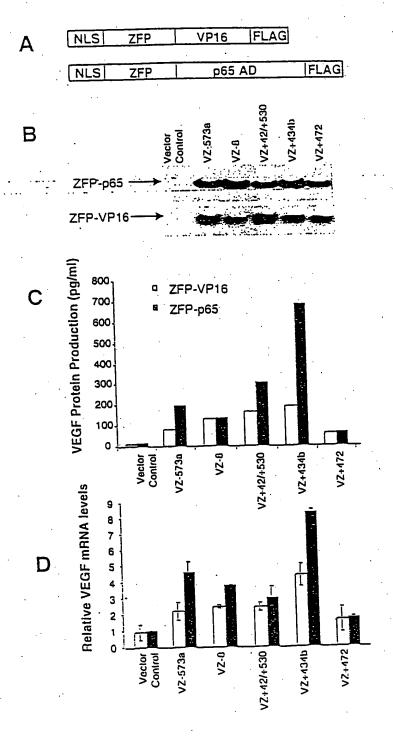


Figure 6

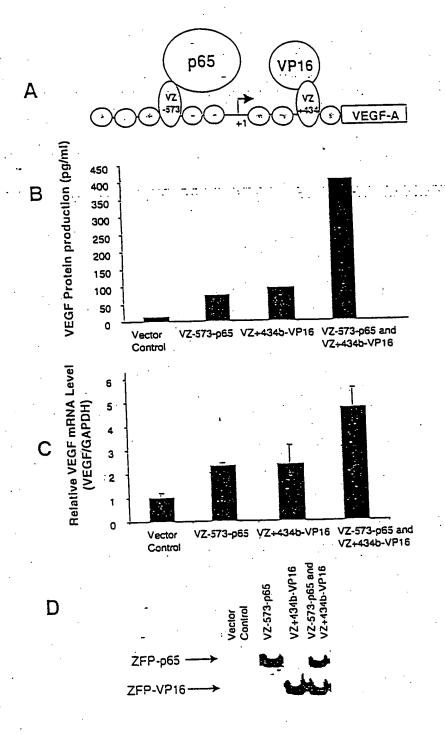
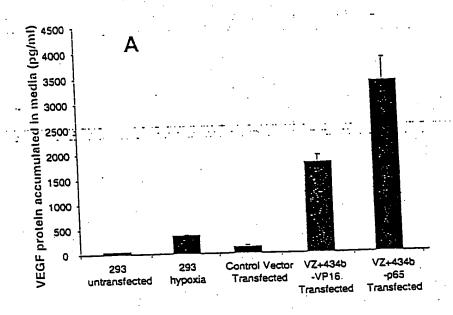


Figure 7



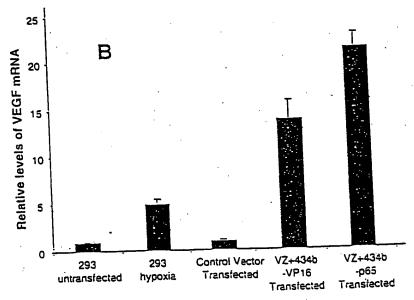
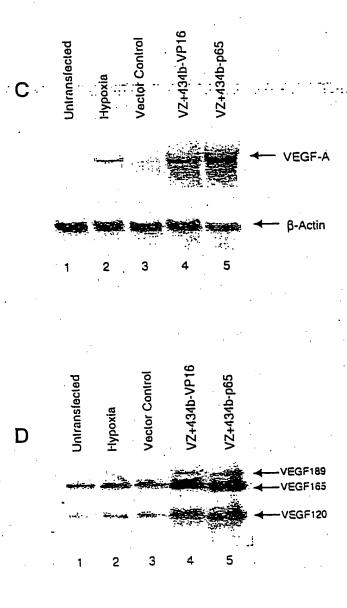


Figure 7

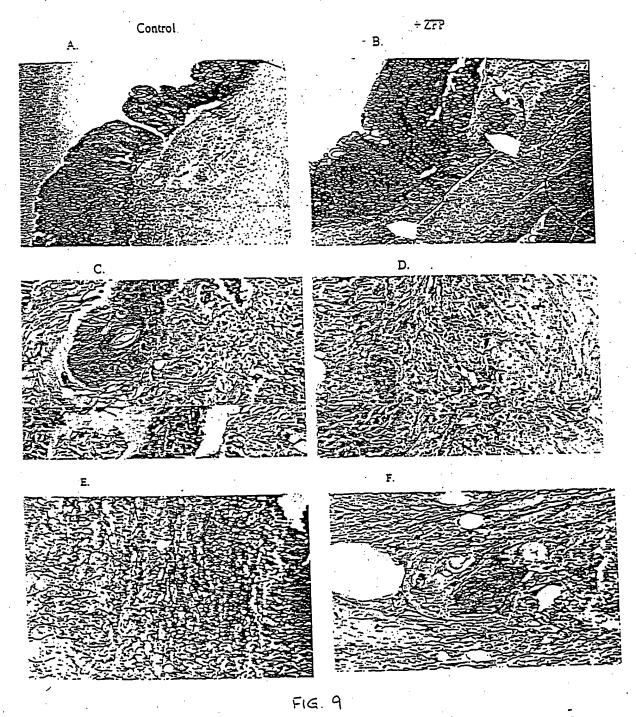


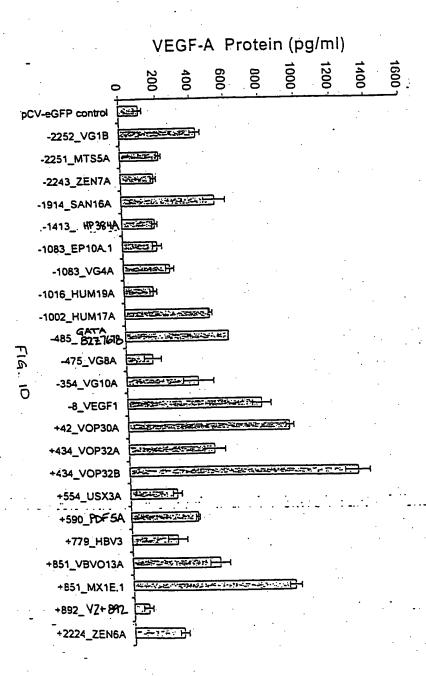
+ZFP Control

Monomer

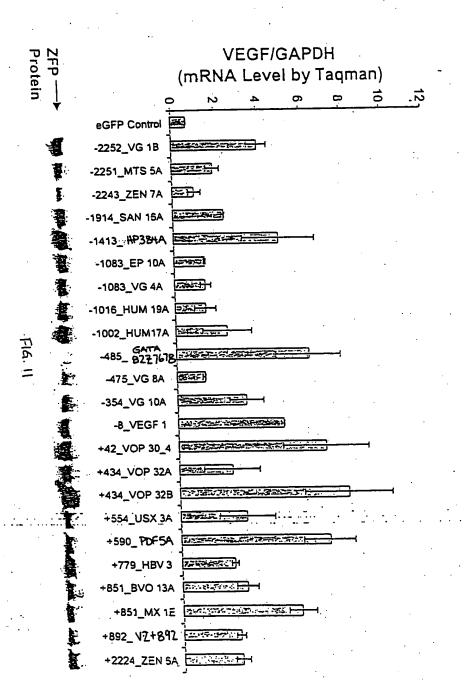
Dimer

FIG. 8



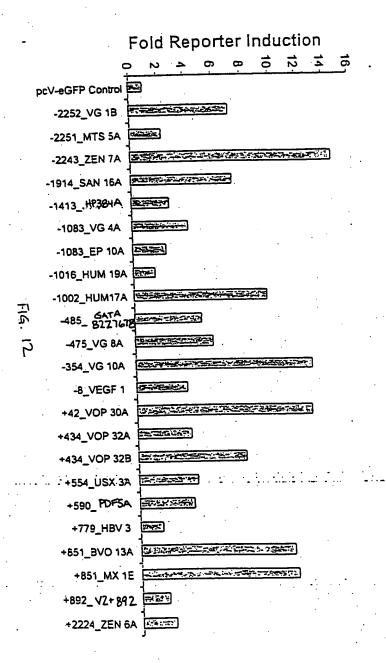


ctivation of Human VEGF-A Gene By ZFPs in 293 Cells (VEGF-A protein production detected by ELISA)

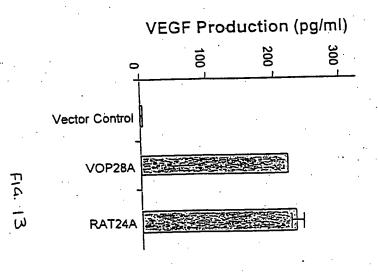


# /EGF-A mRNA Detected by Taqman Analysis Activation of Human VEGF-A Gene by ZFPs

# vation of Human VEGF-A Gene Promoter by ZFPs (Activation of VEGF Reporter in 293 cells)



VEGF Activation By ZFP VOP28A and RAT24A in 293 Cells (VEGF protein detected by ELISA)



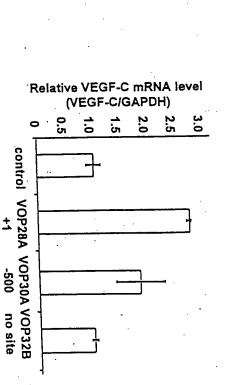


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VEGF-C mRNA Activation by ZFPs

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VEGF-A mRNA Activation by ZFPs

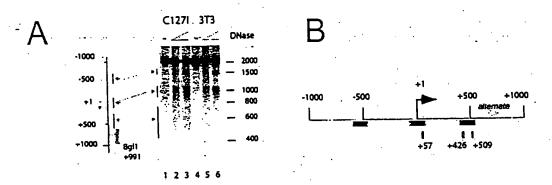


Relative VEGF-A mRNA level (VEGF-A/GAPDH)

control

VOP28A VOP30A VOP32B -500 +1/+5ω +500

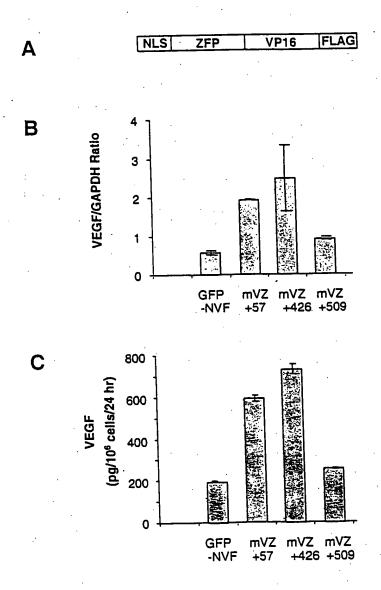
## Figure 15



ZFP Name	Target		Finger	
	Sequence 1-1	Subsites 5-7	Designs 4123448	
±VZ+57	TGAGCGGGGGCAGCGGAGG	CACC CCAC CCAC CCAC CCAC CCAC CCAC CACC	QSGHLTX RSCE_SR RSDELTR QSGSLTR RSCELOR RSCMLAR	
±V2⊷16	ccccctcAC=	6003 6003 6003	rschlir Tschlyr Drsmltr	
mV2+309	GCTGGGGGC4	6664 6664 6614	OSSELTR RSDELTR DRSHLTR	

D	ZFP	Target	Gel Shift		
	mVZ+S7	TGAGCGGCGGCAGCGGAGC		Bound Free	0.031
	mVZ+426	GGGGGTGACc		Bound Free	<0.01
	mVZ+\$09	GCTGGGGGCg		Bound Free	<0.01
	SP1	GGGGCGGGG		Bound Free	0.053

Figure 16



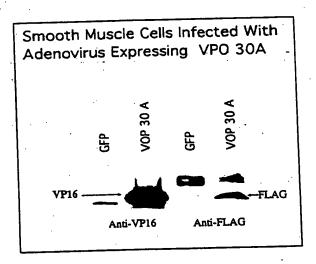


FIG. ITA

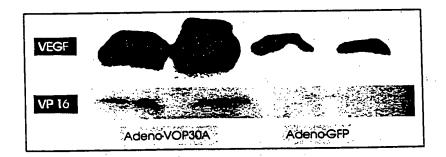


FIG. 17B

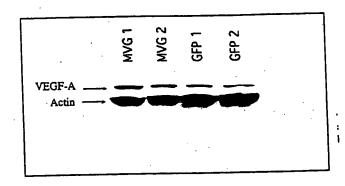


FIG. 17C

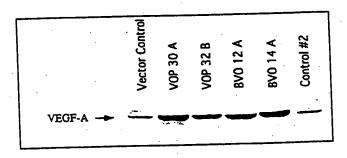


FIG. I7D

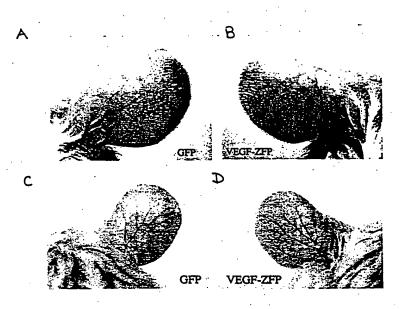


FIG. 18

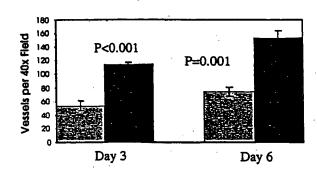
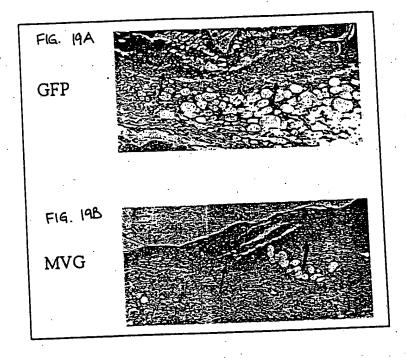


FIG. 18 E



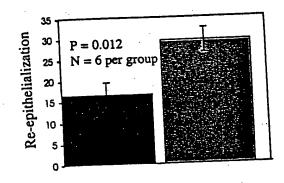


FIG. 19C



VEGF-ZFP (MVG)

FIG. 20A



GFP Control

FIG. 20B

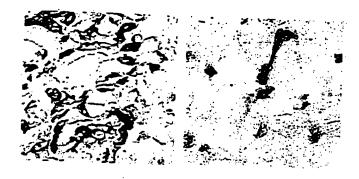


FIG. 21 A

F16. 21 B

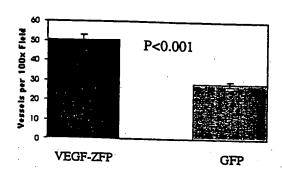


FIG. 21C